

11100	Singh, Sharat Matray, Tracy Chenna, Ahmed	
-:120:- Prok	Sets of Generalized Target-Binding e-tag bes	
+(130)+	w. 25-0033.24	
	Not Yet Assigned Filed Herewith	
	UC 097698,846 1700-10-27	
	UD 09/684,386 .0000-10-04	
	UC 09/602,586 . 00-16-21	
	NO 097861,579 U000-04-28	
	M.: 09/303,029 1499-04-30	
(160 -	18	
-(1 %) -	FistSEQ for Windows Version 4.0	
#210 + #211 + #211 + #2113 +	16	
0.1.10 · 0.1.13 ·	oligonusleotide	
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<pre><pre><pre><pre><pre><pre><pre><pre></pre></pre></pre></pre></pre></pre></pre></pre>	16	
0.1.10 × (2.25 ×	oligonucleotide	
- 40) Jaguu	aggtt tggctg	16

-(210% 3

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:211 - 33
 SIMIN DNA
 +2:13 Artificial Sequence
 -1320 -
 HABBY oligonucleotide
sudd - misc feature
 ·:322 · (32) ... (22)
 4223 * 3' nucleotide linked to tetramethyl rhodamine
+14000-3
                                                                                                                                                                                                                                                                                                22
 mbaqdaadda atgatgooog tt
 -0.10 - 4
 40011 + 22
 HILL - DNA
 Antificial Sequence
H22 - oligonucleotide
Hill misc feature
+1...1. + (1)...(1)
 Para Prince Prin
- L.I - misc_feature
+0.05 \cdot (0.2)^{-1} \cdot ... (22)
Fig. 8' nucleotide linked to tetramethyl rhodamine
  4000
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Howardaarda otgatgootg tt
501.1006
-1. 111. 4
HALL PRI
HULLS Artificial Sequence
-121211
HILE peptide linker
 414.0004 (5)
  Lys Lys Ala Ala
 412100 E
 \pm 1.1111 \pm 4
 · D D PF.T
 -213 Artificial Sequence
*11.2 OD*
 HLIB: pertide linker
 -:400:- 6
 Lys Lys Lys Ala
   1
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<pre>+C108 7 +C110 4 +C110 PRT +C1130 Artificial Sequence</pre>	
<pre>%M240 </pre>	
H40 M T Lys Lys Lys Lys 1	
+2110-36 +2110-16 +2120-DNA +2130-Artificial Sequence	
FLUGA Flugaria de la composición del composición de la composición de la composición de la composición del composición de la composición del composición de la composición del composición del composición del composición del composición del composición del composici	
(4)pp) в фассыдныма tagagaggam atgta	25
+V160 + W110 L7 T170 DNA +V130 Artificial Sequence	
-thto: -thto://wiigonucleotide	
qaaqqaqaag gaagagttgg tattato	27
+0100-10 +0110-1 +0110-0HA +0130-Artificial Sequence	
1730) N.13- Sigonucleotide	
-4000-10 tigggarbag atotgigata g	21
+C100+11 +C110+C7 +C11C+DNA +C113+Artificial Sequence	
<pre><ihito> <li>&lt;2230 oligonucleotide</li></ihito></pre>	
kuunna li patumaggta tobaaaagga gagtota	27
-:210> 12	

K211 × 27 K212 × DNA K213 × Artificial Sequence	
-Q200- -Q203Oligonucleotide	
-04008-12 -oggtatatag ttottootoa tgotatt	27
::10 + 13 -:211 + 20 -:212 + 5MA -::13 + Artificial Sequence	
::::::::::::::::::::::::::::::::::::::	
-:400 - 13 quaugatott ogoottactg	20
-0016 + 14 -0011 + 10 -0010 + DNA -0013 + Artificial Sequence	
-0.110 - -0.113 - probe	
<pre>%Cll = misc_feature %Cll = (1)(1) %Cll = (-tagl0s modification to the 5' nucleotide</pre>	
$<\!(400+14)$ . The satisfies the state of the same space of the satisfies the satisfies and the satisfies the satisfies and the satisfies a satisfies	32
0010 - 15 -0011 - 32 -0010 - DNA -0013 - Artificial Sequence	
Runiu R Runius - probe	
-M21 misc_feature +M220 (1)(1) +M230 +-tagl0as modification to the 5' nucleotide	
04000-15 totitgiata otgototaaa aagaaaatgg aa	32
<pre>F210:- 16 F211:- 08 F212:- DNA F213:- Artificial Sequence</pre>	

+:220>

-MM30- probe	
<pre>chill misc_feature cd2.3 (1)(1) child e-taglls modification to the 5' nucleotide</pre>	
ंपिकाः 16 बब्द : ccago atagatgtgg atagottg	28
<pre>AULIAN 17 AULIAN 28 AULIAN DNA Autificial Sequence</pre>	
-112 t -112 t probe	
<pre>HEAD masc_feature HEAD = (1)(1) HEAD = taglias modification to the 5' nucleotide</pre>	
-:400:- 17 	28
RELIGION 18 RELIGIONA RELI	
-am ar -am r probe	
<pre>HIMTH misc_feature HIMTH (1+(1) HIMTH e-tagl3as modification to the 5' nucleotide</pre>	
<pre>-:400&gt; 18 aactgettgt ggecatgget tag</pre>	23